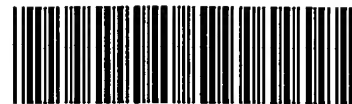# RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/530,844 A

Source: PCT

Date Processed by STIC: 3/10/06

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RAW SEQUENCE LISTING                      DATE: 03/10/2006
PATENT APPLICATION:  US/10/530,844A       TIME: 10:34:23

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Output Set: N:\CRF4\03092006\J530844A.raw
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  6 <120> TITLE OF INVENTION: Feedback-resistant homoserine-Transsuccinylases
having a modified C-
  7         terminus
 10 <130> FILE REFERENCE: Co10221
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C--> 13 <141> CURRENT FILING DATE: 2005-04-08
 15 <160> NUMBER OF SEQ ID NOS: 12
 17 <170> SOFTWARE: PatentIn Ver. 2.0
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 930
 21 <212> TYPE: DNA
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 24 <220> FEATURE:
 25 <221> NAME/KEY: CDS
 26 <222> LOCATION: (1)..(930)
 28 <300> PUBLICATION INFORMATION:
 29 <301> AUTHORs: Blattner, F. R.
 30 <302> TITLE: The complete genome sequence of Escherichia coli K-12.
 31 <303> JOURNAL: Science
 32 <304> VOLUME: 277
 33 <305> ISSUE: 533
 34 <306> PAGES: 1453-1474
 35 <307> DATE: 1997
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 40   1               5                  10                  15
 42  gaa gaa aac gtc ttt gtg atg aca act tct cgt gcg tct ggt cag gaa   96
 43  Glu Glu Asn Val Phe Val Met Thr Thr Ser Arg Ala Ser Gly Gln Glu
 44                  20                  25                  30
 48  att cgt cca ctt aag gtt ctg atc ctt aac ctg atg ccg aag aag att  144
 49  Ile Arg Pro Leu Lys Val Leu Ile Leu Asn Leu Met Pro Lys Lys Ile
 50                  35                  40                  45
 52  gaa act gaa aat cag ttt ctg cgc ctg ctt tca aac tca cct ttg cag  192
 53  Glu Thr Glu Asn Gln Phe Leu Arg Leu Leu Ser Asn Ser Pro Leu Gln
 54          50                  55                  60
 56  gtc gat att cag ctg ttg cgc atc gat tcc cgt gaa tcg cgc aac acg  240
 57  Val Asp Ile Gln Leu Leu Arg Ile Asp Ser Arg Glu Ser Arg Asn Thr
 58  65                  70                  75                  80
 60  ccc gca gag cat ctg aac aac ttc tac tgt aac ttt gaa gat att cag  288
 61  Pro Ala Glu His Leu Asn Asn Phe Tyr Cys Asn Phe Glu Asp Ile Gln
 62                  85                  90                  95
 64  gat cag aac ttt gac ggt ttg att gta act ggt gcg ccg ctg ggc ctg  336
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69   Val Glu Phe Asn Asp Val Ala Tyr Trp Pro Gln Ile Lys Gln Val Leu
70           115                     120                 125
72   gag tgg tcg aaa gat cac gtc acc tcg acg ctg ttt gtc tgc tgg gcg      432
73   Glu Trp Ser Lys Asp His Val Thr Ser Thr Leu Phe Val Cys Trp Ala
74           130                     135                 140
76   gta cag gcc gcg ctc aat atc ctc tac ggc att cct aag caa act cgc      480
77   Val Gln Ala Ala Leu Asn Ile Leu Tyr Gly Ile Pro Lys Gln Thr Arg
78   145                 150                     155                 160
80   acc gaa aaa ctc tct ggc gtt tac gag cat cat att ctc cat cct cat      528
81   Thr Glu Lys Leu Ser Gly Val Tyr Glu His His Ile Leu His Pro His
82                   165                     170                 175
84   gcg ctt ctg acg cgt ggc ttt gat gat tca ttc ctg gca ccg cat tcg      576
85   Ala Leu Leu Thr Arg Gly Phe Asp Asp Ser Phe Leu Ala Pro His Ser
86               180                     185                 190
88   cgc tat gct gac ttt ccg gca gcg ttg att cgt gat tac acc gat ctg      624
89   Arg Tyr Ala Asp Phe Pro Ala Ala Leu Ile Arg Asp Tyr Thr Asp Leu
90           195                     200                 205
92   gaa att ctg gca gag acg gaa gaa ggg gat gca tat ctg ttt gcc agt      672
95   Glu Ile Leu Ala Glu Thr Glu Glu Gly Asp Ala Tyr Leu Phe Ala Ser
96           210                     215                 220
98   aaa gat aag cgc att gcc ttt gtg acg ggc cat ccc gaa tat gat gcg      720
99   Lys Asp Lys Arg Ile Ala Phe Val Thr Gly His Pro Glu Tyr Asp Ala
100  225                     230                     235                 240
102  caa acg ctg gcg cag gaa ttt ttc cgc gat gtg gaa gcc gga cta gac      768
103  Gln Thr Leu Ala Gln Glu Phe Phe Arg Asp Val Glu Ala Gly Leu Asp
104                   245                     250                 255
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107  Pro Asp Val Pro Tyr Asn Tyr Phe Pro His Asn Asp Pro Gln Asn Thr
108                   260                     265                 270
110  ccg cga gcg agc tgg cgt agt cac ggt aat tta ctg ttt acc aac tgg      864
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112           275                     280                 285
114  ctc aac tat tac gtc tac cag atc acg cca tac gat cta cgg cac atg      912
115  Leu Asn Tyr Tyr Val Tyr Gln Ile Thr Pro Tyr Asp Leu Arg His Met
116           290                     295                 300
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Input Set : A:\PTO.ss.txt
Output Set: N:\CRF4\03092006\J530844A.raw

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143    Val Asp Ile Gln Leu Leu Arg Ile Asp Ser Arg Glu Ser Arg Asn Thr
144      65                70                75                80
146    Pro Ala Glu His Leu Asn Asn Phe Tyr Cys Asn Phe Glu Asp Ile Gln
147                85                90                95
149    Asp Gln Asn Phe Asp Gly Leu Ile Val Thr Gly Ala Pro Leu Gly Leu
150             100               105               110
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153          115               120               125
155    Glu Trp Ser Lys Asp His Val Thr Ser Thr Leu Phe Val Cys Trp Ala
156          130               135               140
158    Val Gln Ala Ala Leu Asn Ile Leu Tyr Gly Ile Pro Lys Gln Thr Arg
159    145               150               155               160
161    Thr Glu Lys Leu Ser Gly Val Tyr Glu His His Ile Leu His Pro His
162               165               170               175
164    Ala Leu Leu Thr Arg Gly Phe Asp Asp Ser Phe Leu Ala Pro His Ser
165               180               185               190
167    Arg Tyr Ala Asp Phe Pro Ala Ala Leu Ile Arg Asp Tyr Thr Asp Leu
168             195               200               205
170    Glu Ile Leu Ala Glu Thr Glu Glu Gly Asp Ala Tyr Leu Phe Ala Ser
171          210               215               220
173    Lys Asp Lys Arg Ile Ala Phe Val Thr Gly His Pro Glu Tyr Asp Ala
174    225               230               235               240
176    Gln Thr Leu Ala Gln Glu Phe Phe Arg Asp Val Glu Ala Gly Leu Asp
177               245               250               255
179    Pro Asp Val Pro Tyr Asn Tyr Phe Pro His Asn Asp Pro Gln Asn Thr
180             260               265               270
182    Pro Arg Ala Ser Trp Arg Ser His Gly Asn Leu Leu Phe Thr Asn Trp
183          275               280               285
185    Leu Asn Tyr Tyr Val Tyr Gln Ile Thr Pro Tyr Asp Leu Arg His Met
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268      gene
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299   gene
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308 Gly Ala
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336   gene
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343   20   25   30
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 03/10/2006
PATENT APPLICATION: US/10/530,844A   TIME: 10:34:24

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Output Set: **N:\CRF4\03092006\J530844A.raw**

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VERIFICATION SUMMARY   DATE: 03/10/2006
PATENT APPLICATION: US/10/530,844A   TIME: 10:34:24

Input Set : A:\PTO.ss.txt
Output Set: N:\CRF4\03092006\J530844A.raw

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